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By: Nancy Ramos Printed: Nancy Ramos

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of: Bandman et al.

Title: HUMAN EXTRACELLULAR MATRIX PROTEINS

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SUBMISSION OF FORMAL DRAWINGS

Sir:

Transmitted herewith are Figure(s) 1A, 1B, 1C, 1D, 1E, 1F, 1G, 2A, 2B, 2C, 2D, 2E, 3A, 3B, 4A, 4B, and 4C, as seventeen (17) sheets of formal drawings for this application. Each sheet of drawing indicates the identifying indicia suggested in 37 CFR Section 1.84(c) on the reverse side of the drawings.

Applicants believe that no fee is due with this paper. However, if the Commissioner determines that a fee is necessary, the Commissioner is hereby authorized to charge any additional fees associated with this communication or credit any overpayment to Deposit Account No. **09-0108**. **A duplicate copy of this communication is enclosed.**

If there are any questions regarding the above, the Examiner is invited to call the undersigned at 650-855-0555.

Respectfully submitted,

INCYTE GENOMICS, INC.

Date: 16th April 2001

Matthew R. Kaser
Matthew R. Kaser, D.Phil.
Reg. No. 44,817
Direct Dial Telephone: (650) 845-4596

3160 Porter Drive
Palo Alto, California 94304
Phone: (650) 855-0555
Fax: (650) 845-4166

[illegible]

FIGURE 1B

819	828	837	846	855	864
TGC ATC AAT	GGC GGC GGC	TAC ACC	TGC TCC	GAC GGA	TAT TGG
C I N	E G G	Y T C	S C C	D G Y	W L
873	882	891	900	909	918
CTG GAA GGC	TGC TTA GAC	ATT GAT	TAT CGC	GGT TAC	TGC CAG
L E G	C L C	I D E	R Y C	Y G C	Q Q
927	936	945	954	963	972
CTC TGT GCG	AAT GTT CCT	TCC TAT	TGT ACA	AAC CCT	GGT TTT
L C A	N V P	S Y S	T C C	N P G	F T
981	990	999	1008	1017	1026
CTC AAT GAG	GAT GGA AGG	TGC CAA	GAT GTG	TGT GCC	ACC GAG
L N E	D G R	C Q D	N V D	C A T	E N
1035	1044	1053	1062	1071	1080
CCC TGC GTG	CAA ACC TGC	AAC ACC	TGC TCT	ATC TGC	CGC TGT
P C V	Q T C	N T Y	S G C	I C R	C D
1089	1098	1107	1116	1125	1134
CCA GGA TAT	GAA CTT GAG	GAT GGC	TGC CAT	GAT ATG	GAG TGC
P G Y	E L E	D G V	H C S	M D E	C C
1143	1152	1161	1170	1179	1188
AGC TTC TCT	GAG TTC CTC	CAA CAT	TGT GTG	CAG CCC	GGC ACA
S F S	E F L	O C	V C	Q P G	T T

FIGURE 1C

1197 1206 1215 1224 1233 1242
 TTC TGC TCC TGC CCT CCA GGC TAC ATC CTG CTG GAT GAC AAC CGA AGC TGC CAA
 F C S C P P G Y I L L D D N R S C Q

1251 1260 1269 1278 1287 1296
 GAC ATC AAC GAA TGT GAG CAC AGG AAC CAC ACG TGC AAC CTG CAG CAG ACG TGC
 D I N E C E H R N H T C N L Q Q T C

1305 1314 1323 1332 1341 1350
 TAC AAT TTA CAA GGC GGC TTC AAA TGC ATC GAC CCC ATC CGC TGT GAG GAG CCT
 Y N L Q G G F K C I D P I R C E E P

1359 1368 1377 1386 1395 1404
 TAT CTG AGG ATC AGT GAT AAC CGC TGT ATG TGT CCT GCT GAG AAC CCT GGC TGC
 Y L R I S D N R C M C P A E N P G C

1413 1422 1431 1440 1449 1458
 AGA GAC CAG CCC TTT ACC ATC TTG TAC CGG GAC ATG GAC GTG GTG TCA GGA CGC
 R D Q P F T I L Y R D M D V V S G R

1467 1476 1485 1494 1503 1512
 TCC GTT CCC GCT GAC ATC TTC CAA ATG CAA GCC ACG ACC CGC TAC CCT GGC GCC
 S V P A D I F Q M Q A T T R Y P G A

1521 1530 1539 1548 1557 1566
 TAT TAC ATT TTC CAG ATC AAA TCT GGC AAT GAG GGC AGA GAA TTT TAC ATG CGG
 Y Y I F Q I K S G N E G R E F Y M R

FIGURE 1D

1575 1584 1593 1602 1611 1620
 CAA ACG GGC CCC ATC AGT GGC ACC CTG GTG ATG ACA CGC CCC ATC AAA GGG CCC
 Q T G P I S A T L V M T R P I K G P

1629 1638 1647 1656 1665 1674
 CGG GAA ATC CAG CTG GAC TTG GAA ATG ATC ACT GTC AAC ACT GTC ATC AAC TTC
 R E I Q L D L E M I T V N T V I N F

1683 1692 1701 1710 1719 1728
 AGA GGC AGC TCC GTG ATC CGA CTG CGG ATA TAT GTG TCG CAG TAC CCA TTC TGA
 R G S S V I R L R I Y V S Q Y P F

1737 1746 1755 1764 1773 1782
 GCC TCG GGC TGG AGC CTC CGA CGC TGC CTC TCA TTG GCA CCA AGG GAC AGG AGA
 AGA GAG GAA ATA ACA GAG AGA ATG AGA GCG ACA CAG ACG TTA GGC ATT TCC TGC

1791 1800 1809 1818 1827 1836
 AGA GAG GAA ATA ACA GAG AGA ATG AGA GCG ACA CAG ACG TTA GGC ATT TCC TGC
 1845 1854 1863 1872 1881 1890
 TGA ACG TTT CCC CGA AGA GTC AGC CCC GAC TTC CTG ACT CTC ACC TGT ACT ATT

1899 1908 1917 1926 1935 1944
 GCA GAC CTG TCA CCC TGC AGG ACT TGC CAC CCC CAG TTC CTA TGA CAC AGT TAT
 1953 1962 1971 1980 1989 1998
 CAA AAA GTA TTA TCA TTG CTC CCC TGA TAG AAG ATT GTT GGT GAA TTT TCA AGG

FIGURE 1E

2007	2016	2025	2034	2043	2052
CCT TCA GTT TAT TTC CAC TAT TTT CAA AGA AAA TAG ATT AGG TTT GCG GGC GTC					
2061	2070	2079	2088	2097	2106
TGA GTC TAT GTT CAA AGA CTG TGA ACA GCT TGC TGT CAC TTC TTC ACC TCT TCC					
2115	2124	2133	2142	2151	2160
ACT CCT TCT CTC ACT GTG TTA CTG CTT TGC AAA GAC CCG GGA GCT GGC GGC GAA					
2169	2178	2187	2196	2205	2214
CCC TGG GAG TAG CTA GTT TGC TTT TTG CGT ACA CAG AGA AGG CTA TGT AAA CAA					
2223	2232	2241	2250	2259	2268
ACC ACA GCA GGA TCG AAG GGT TTT TAG AGA ATG TGT TTC AAA ACC ATG CCT GGT					
2277	2286	2295	2304	2313	2322
ATT TTC AAC CAT AAA AGA AGT TTC AGT TGT CCT TAA ATT TGT ATA ACG GTT TAA					
2331	2340	2349	2358	2367	2376
TTC TGT CTT GTT CAT TTT GAG TAT TTT TAA AAA ATA TGT CGT AGA ATT CCT TCG					
2385	2394	2403	2412	2421	2430
AAA GGC CTT CAG ACA CAT GCT ATG TTC TGT CTT CCC AAA CCC AGT CTC CTC TCC					
2439	2448	2457	2466	2475	2484
ATT TTA GCC CAG TGT TTT CTT TGA GGA CCC CTT AAT CTT GCT TTC TTT AGA ATT					

FIGURE 1F

2493 2502 2511 2520 2529 2538
TTT ACC CAA TTG GAT TGG AAT GCA GAG GTC TCC AAA CTG ATT AAA TAT TTG AAG
2547
AGA AAA AAA AAA

FIGURE 1G

1000 1000 1000 1000 1000 1000 1000 1000 1000 1000
 1000 1000 1000 1000 1000 1000 1000 1000 1000 1000
 1000 1000 1000 1000 1000 1000 1000 1000 1000 1000

9	18	27	36	45	54
TG GGT GCA AGC TCA CAA CCG TAA CAG CCA CCA GAC AAG CTT CAG TGG CCG GCC					
63	72	81	90	99	108
CTT CAC ATC CAG ACT TGC CTG AGA GGA CCC ACC TCT GAG TGT CCA GTG GTC AGT					
117	126	135	144	153	162
TGC CCC AGG ATG GGG ACC ACA GCC AGA GCA GCC TTG GTC TTG ACC TAT TTG GCT					
M G T T A R A A A L V L Y L A					
171	180	189	198	207	216
GTT GCT TCT GCT GCC TCT GAG GGA GGC TTC ACG GCT ACA GGA CAG AGG CAG CTG					
V A S A A S E G G F T A T G Q R Q L					
225	234	243	252	261	270
AGG CCA GAG CAC TTT CAA GAA GTT GGC TAC GCA GCT CCC CCC CCA CCC CTA					
R P E H F Q E V G Y A A P P S P L					
279	288	297	306	315	324
TCC CGA AGC CTC CCC ATG GAT CAC CCT GAC TCC TCT CAG CAT GGC CCT CCC TTT					
S R S L P M D H P D S S Q H G P P F					
333	342	351	360	369	378
GAG GGA CAG AGT CAA GTG CAG CCC CCT CCC TCT CAG GAG GCC ACC CCT CTC CAA					
E G Q S Q V Q P P P S Q Q E A T P L Q					

FIGURE 2A

387	CAG GAA AAG CTG CTA CCT GCC CAA CTC CCT GCT GAA AAG GAA GTG GGT CCC CCT	405	414	423	432
Q E K L L P A Q L P A E K E V G P P					
441	CTC CCT CAG GAA GCT GTC CCC CTC CAA AAA GAG CTG CCC TCT CTC CAG CAC CCC	459	468	477	486
L P Q E A V P L Q K E L P S L Q H P					
495	AAT GAA CAG AAG GAA GGA ATG CCA GCT CCA TTT GGG GAC CAG AGC CAT CCA GAA	513	522	531	540
N E Q K E G M P A P F G D Q S H P E					
549	CCT GAG TCC TGG AAT GCA GCC CAG CAC TGC CAA CAG GAC CGG TCC CAA GGG GGC	567	576	585	594
P E S W N A A Q H C Q Q D R S Q G G					
603	TGG GGC CAC CGG CTG GAT GGC TTC CCC CCT GGG CGG CCT TCT CCA GAC AAT CTG	621	630	639	648
W G H R L D G F P P G R P S P D N L					
657	AAC CAA ATC TGC CTT CCT AAC CGT CAG CAT GTG GTA TAT GGT CCC TGG AAC CTA	675	684	693	702
N Q I C L P N R Q H V V Y G P W N L					
711	CCA CAG TCC AGC TAC TCC CAC CTC ACT CGC CAG GGT GAG ACC CTC AAT TTC CTG	729	738	747	756
P Q S S Y S H L T R Q G E T L N F L					

FIGURE 2B

1143 1144 1145 1146 1147 1148 1149 1150 1151 1152 1153 1154 1155 1156 1157 1158 1159 1160 1161 1162 1163 1164 1165 1166 1167 1168 1169 1170 1171 1172 1173 1174 1175 1176 1177 1178 1179 1180 1181 1182 1183 1184 1185 1186 1187 1188 1189 1190 1191 1192 1193 1194 1195 1196 1197 1198 1199 1200 1201 1202 1203 1204 1205 1206 1207 1208 1209 1210 1211 1212 1213 1214 1215 1216 1217 1218 1219 1220 1221 1222 1223 1224 1225 1226 1227 1228 1229 1230 1231 1232 1233 1234 1235 1236 1237 1238 1239 1240 1241 1242 1243 1244 1245 1246 1247 1248 1249 1250 1251 1252 1253 1254 1255 1256 1257 1258 1259 1260 1261 1262 1263 1264 1265 1266 1267 1268 1269 1270 1271 1272 1273 1274 1275 1276 1277 1278 1279 1280 1281 1282 1283 1284 1285 1286 1287 1288 1289 1290 1291 1292 1293 1294 1295 1296 1297 1298 1299 1300 1301 1302 1303 1304 1305 1306 1307 1308 1309 1310 1311 1312 1313 1314 1315 1316 1317 1318 1319 1320 1321 1322 1323 1324 1325 1326 1327 1328 1329 1330 1331 1332 1333 1334 1335 1336 1337 1338 1339 1340 1341 1342 1343 1344 1345 1346 1347 1348 1349 1350 1351 1352 1353 1354 1355 1356 1357 1358 1359 1360 1361 1362 1363 1364 1365 1366 1367 1368 1369 1370 1371 1372 1373 1374 1375 1376 1377 1378 1379 1380 1381 1382 1383 1384 1385 1386 1387 1388 1389 1390 1391 1392 1393 1394 1395 1396 1397 1398 1399 1400 1401 1402 1403 1404 1405 1406 1407 1408 1409 1410 1411 1412 1413 1414 1415 1416 1417 1418 1419 1420 1421 1422 1423 1424 1425 1426 1427 1428 1429 1430 1431 1432 1433 1434 1435 1436 1437 1438 1439 1440 1441 1442 1443 1444 1445 1446 1447 1448 1449 1450 1451 1452 1453 1454 1455 1456 1457 1458 1459 1460 1461 1462 1463 1464 1465 1466 1467 1468 1469 1470 1471 1472 1473 1474 1475 1476 1477 1478 1479 1480 1481 1482 1483 1484 1485 1486 1487 1488 1489 1490 1491 1492 1493 1494 1495 1496 1497 1498 1499 1500 1501 1502 1503 1504 1505 1506 1507 1508 1509 1510 1511 1512 1513 1514 1515 1516 1517 1518 1519 1520 1521 1522 1523 1524 1525 1526 1527 1528 1529 1530 1531 1532 1533 1534 1535 1536 1537 1538 1539 1540 1541 1542 1543 1544 1545 1546 1547 1548 1549 1550 1551 1552 1553 1554 1555 1556 1557 1558 1559 1560 1561 1562 1563 1564 1565 1566 1567 1568 1569 1570 1571 1572 1573 1574 1575 1576 1577 1578 1579 1580 1581 1582 1583 1584 1585 1586 1587 1588 1589 1590 1591 1592 1593 1594 1595 1596 1597 1598 1599 1600

1143	1152	1161	1170	1179	1188
CAG CTG GAG AGG GAG TTC CAG CGC TGC TGC CGC CAG GGG AAC AAT CAC ACC TGT					
Q L E R E F Q R C C R Q G N N H T C					
1197	1206	1215	1224	1233	1242
ACA TGG AAG GCC TGG GAG GAT ACC CTT GAC AAA TAC TGT GAC CGG GAG TAT GCT					
T W K A A W E D T L D K Y C D R E Y A					
1251	1260	1269	1278	1287	1296
GTG AAG ACC CAC CAC CAC TTG TGT TGC CGC CAC CCT CCC AGC CCT ACT CGG GAT					
V K T H H H L C C R H P P S P T R D					
1305	1314	1323	1332	1341	1350
GAG TGC TTT GCC CGT CGG GCT CCT TAC CCC AAC TAT GAC CGG GAC ATC TTG ACC					
E C F A R R A P Y P N Y D R D I L T					
1359	1368	1377	1386	1395	1404
ATT GAC ATC GGT CGA GTC ACC CCC AAC CTC ATG GGC CAC CTC TGT GGA AAC CAA					
I D I G R V T P N L M G H L C G N Q					
1413	1422	1431	1440	1449	1458
AGA GTT CTC ACC AAG CAT AAA CAT ATT CCT GGG CTG ATC CAC AAC ATG ACT GCC					
R V L T K H K H I P G L I H N M T A					
1467	1476	1485	1494	1503	1512
CGC TGC TGT GAC CTG CCA TTT CCA GAA CAG GCC TGC TGT GCA GAG GAG GAG AAA					
R C C D L P F P E Q A C C A E E E K					

FIGURE 2D

1521	TTA ACC TTC ATC AAT GAT CTG TGT GGT CCC CGA CGT AAC ATC TGG CGA GAC CCT	1530	1539	1548	1557	1566
L T F I N D L C G P R R R I W R D P						
1575	GCC CTC TGC TGT TAC CTG AGT CCT GGG GAT GAA CAG GTC AAC TGC TTC AAC ATC	1584	1593	1602	1611	1620
A L C C C Y L S P G D E Q V N C F N I						
1629	AAT TAT CTG AGG AAC GTG GCT CTA GTG TCT GGA GAC ACT GAG AAC GCC AAC GGC	1638	1647	1656	1665	1674
N Y L R N V A L V S G D T E N A K G						
1683	CAG GGG GAG CAG GGC TCA ACT GGA GGA ACA AAT ATC AGC TCC ACC TCT GAG CCC	1692	1701	1710	1719	1728
Q G E Q Q G S T G G T N I S S T S E P						
1737	AAG GAA GAA TGA GTC ACC CCA GAG CCC TAG AGG GTC AGA TGG GGG GAA CCC CAC	1746	1755	1764	1773	1782
K E E						
1791	CCT GCC CCA CCC ATC TGA ACA CTC ATT ACA CTA AAC ACC TCT TGG ATT TGG TGT	1800	1809	1818	1827	1836
1845	CCT CAT TGT CTA TCT AAT GTC TCA CCC GCA GTG TTT TAA GTG GAT CTT GGT GCC	1854	1863	1872	1881	1890
1899	CTG GCC CAG G					

FIGURE 2E

1	M P G I K R I L T V T I L A L C L P S P	45517
1	M A T S G V L P G G G F V A S A A A V A	GI 458228
21	G N A Q A Q C T N G F D L D R Q S G Q C	45517
21	G P E M Q T G R N N F V I R R N P A D -	GI 458228
41	L D I D E C R T I P E A C R G D M M C V	45517
40	- - - - -	GI 458228
61	N Q N G G Y L C I P R T N P V Y R G P Y	45517
40	- - - - -	GI 458228
81	S N P Y S T P Y S G P Y P A A A P P L S	45517
40	- - - - - P Q R	GI 458228
101	A P N Y P T I S R P L I C R F G Y Q M D	45517
43	I P S N P - - S H R I Q C A A G Y E Q S	GI 458228
121	E S N Q C V D V D E C A T D S H Q C N P	45517
61	E H N V C Q D I D E C T A G T H N C R A	GI 458228
141	T Q I C I N T E G G Y T C S C T D G Y W	45517
81	D Q V C I N L R G S F A C Q C P P G Y Q	GI 458228
161	L L E G Q C L D I D E C R Y G - Y C Q Q	45517
101	K R G E Q C V D I D E C T I P P Y C H Q	GI 458228
180	L C A N V P G S Y S C T C N P G F T L N	45517
121	R C V N T P G S F Y C Q C S P G F Q L A	GI 458228
200	E D G R S C Q D V N E C A T E N P C V Q	45517
141	A N N Y T C V D I N E C D A S N Q C A Q	GI 458228
220	T C V N T Y G S F I C R C D P G Y E L E	45517
161	Q C Y N I L G S F I C Q C N Q G Y E L S	GI 458228

FIGURE 3A

240 E D G V H C S D M D E C S F S E F L C Q 45517
 181 S D R L N C E D I D E C R T S S Y L C Q GI 458228

260 H E C V N Q P G T Y F C S C P P G Y I L 45517
 201 Y Q C V N E P G K F S C M C P Q G Y Q V GI 458228

280 L D D N R S C Q D I N E C E H R N H T C 45517
 221 V R - S R T C Q D I N E C E T T N E - C GI 458228

300 N L Q Q T C Y N L Q G G F K C I D P I R 45517
 239 R E D E M C W N Y H G G F R C Y P R N P GI 458228

320 C E E P Y L R I S D N R C M C P A E N P 45517
 259 C Q D P Y I L T P E N R C V C P V S N A GI 458228

340 G C R D Q P F T I L Y R D M D V V S G R 45517
 279 M C R E L P Q S I V Y K Y M S I R S D R GI 458228

360 S V P A D I F Q M Q A T T R Y P G A Y Y 45517
 299 S V P S D I F Q I Q A T T I Y A N T I N GI 458228

380 I F Q I K S G N E G R E F Y M R Q T G P 45517
 319 T F R I K S G N E N G E F Y L R Q T S P GI 458228

400 I S A T L V M T R P I K G P R E I Q L D 45517
 339 V S A M L V L V K S L S G P R E H I V D GI 458228

420 L E M I T V N T V I N F R G S S V I R L 45517
 359 L E M L T V S S I G T F R T S S V L R L GI 458228

440 R I Y V S Q Y P F 45517
 379 T I I V G P F S F GI 458228

FIGURE 3B

1	M G T T A R A A L V L T Y L A V A S A A	1621777
1	M G T V S R A A L I L A C L A L A S A A	GI 496120
21	S E G G F T A T G Q R Q L R P E - - - -	1621777
21	S E G A F K A S D Q R E M T P E R L F Q	GI 496120
37	H F Q E V G Y A A P P S P P L S R S L P	1621777
41	H L H E V G Y A A P P S L P Q T R R L R	GI 496120
57	M D H P D S S Q H G P P - F E G Q S Q V	1621777
61	V D H S V T S L H D P P L F E E Q R E V	GI 496120
76	Q P P P S Q E A T P L Q Q E K L L P A Q	1621777
81	Q P P S S P E D I P V Y E E D W P T F L	GI 496120
96	L P A E K E V G P P L P Q E A V P L Q K	1621777
101	N P N V D K A G P A V P Q E A I P L Q K	GI 496120
116	E L P S L Q - - - - - - - - - - H P	1621777
121	E Q P P P Q V H I E Q K E I D P P A Q P	GI 496120
124	N E - - - Q K E G M P A P F G D Q S H P	1621777
141	Q E E I V Q K E V K P H T L A G Q L P P	GI 496120
141	E P E S W N A A Q H C Q Q D R S Q G G W	1621777
161	E P R T W N P A R H C Q Q G R - R G V W	GI 496120
161	G H R L D G F P P G R P S P D N L N Q I	1621777
180	G H R L D G F P P G R P S P D N L K Q I	GI 496120
181	C L P N R Q H V V Y G P W N L P Q S S Y	1621777
200	C L P E R Q H V I Y G P W N L P Q T G Y	GI 496120
201	S H L T R Q G E T L N F L E I G Y S R C	1621777
220	S H L S R Q G E T L N V L E T G Y S R C	GI 496120

FIGURE 4A

221 C H C R S H T N R L E C A K L V W E E A 1621777
240 C P C R S D T N R L D C L K L V W E D A GI 496120

241 M S R F C E A E F S V K T R P H W C C T 1621777
260 M T Q F C E A E F S V K T R P H L C C R GI 496120

261 R Q G E A R F S C F Q E E A P Q P H Y Q 1621777
280 L R G E E R F S C F Q K E A P R P D Y L GI 496120

281 L R A C P S H Q P D I S S G L E L P F P 1621777
300 L R P C P V H Q N G M S S G P Q L P F P GI 496120

301 P G V P T L D N I K N I C H L R R F R S 1621777
320 P G L P T P D N V K N I C L L R R F R A GI 496120

321 V P R N L P A T D P L Q R E L L A L I Q 1621777
340 V P R N L P A T D A I Q R Q L Q A L T R GI 496120

341 L E R E F Q R C C R Q G N N H T C T W K 1621777
360 L E T E F Q R C C R Q G H N H T C T W K GI 496120

361 A W E D T L D K Y C D R E Y A V K T H H 1621777
380 A W E G T L D G Y C E R E L A I K T H P GI 496120

381 H L C C R H P P S P T R D E C F A R R A 1621777
400 H S C C H Y P P S P A R D E C F A H L A GI 496120

401 P Y P N Y D R D I L T I D I G R V T P N 1621777
420 P Y P N Y D R D I L T L D L S R V T P N GI 496120

421 L M G H L C G N Q R V L T K H K H I P G 1621777
440 L M G Q L C G S G R V L S K H K Q I P G GI 496120

441 L I H N M T A R C C D L P F P E Q A C C 1621777
460 L I Q N M T V R C C E L P Y P E Q A C C GI 496120

FIGURE 4B

461	A E E E K L T F I N D L C G P R R N I W	1621777
480	G E E E K L A F I E N L C G P R R N S W	GI 496120
481	R D P A L C C Y L S P G D E Q V N C F N	1621777
500	K D P A L C C D L S P E D K Q I N C F N	GI 496120
501	I N Y L R N V A L V S G D T E N A K G Q	1621777
520	T N Y L R N V A L V A G D T G N A T G L	GI 496120
521	G E Q G S T G G T N I S S T S E P K E E	1621777
540	G E Q G P T R G T D A N P A P G S K E E	GI 496120

FIGURE 4C